Table S2 Oligos for amplifying Arabidopsis DNA

| Oligo name | Oligo sequence | Purpose | Note |
|-----------------|--------------------------|--|--|
| | | For detection of mutations at At1g53430 | |
| At1g53430-F2 | CTgtaagcaaaactaactaaccac | site | |
| | ctcacGTTTAGCATCTTCTGGA | For detection of mutations at At1g53430 | Designated as an open arrow in Fig |
| At1g53430-R2 | CA | site and gene cluster inversions | S5A and B |
| | | For detection of gene cluster inversions | Designated as F2 in Fig. 4A and as a |
| At1g53440-F1 | tagatgatatttttaaccgtgac | and large chromosomal deletions | filled, tailess arrow in Fig. 5A and B |
| | | For detection of mutations at At1g53440 | |
| At1g53440-F2 | tattcggtatcatcaaggtca | site | |
| == | TTCTTAAGCACCATTGGACAct | For detection of mutations at At1g53440 | |
| At1g53440-R2 | ac | site | |
| | | For detection of gene cluster deletions, | Designated as F1 in Fig. 3A and Fig. |
| | | inversions and large chromosomal | 4A, and as an open tailess arrow in |
| At1g53430-F1 | ATTGGTTCCATGAGTGAGC | deletions | Fig. 5A and B |
| | | For detection of gene cluster deletions | Designated as R2 in Fig. 3A and as a |
| At1g53440-R3 | aagggcttctttttttcaag | and inversions | filled arrow in Fig. 5A and B |
| | TTCTTCTCCAACAGCACCGTC | For detection of mutations at At1g70450 | |
| At1g70450-F3 | AG | site and gene cluster deletions | Designated as F1 in Fig. 3B |
| | | For detection of mutations at At1g70450 | Designated as an open arrow in Fig |
| At1g70450-R2 | CACTGGCCTACCCTTCCctgtc | site and gene cluster inversions | S8A and B |
| . 0 | | For detection of gene cluster | Designated as an open arrow in Fig |
| At1g70450-R3 | GGTGACctgcaaaacaagataaat | duplications | S9 |
| | TACTCTGGTCCTGGTGGTTAC | For detection of gene cluster | Designated as a tailess filled arrow |
| At1g70460-F | AAT | duplications | in Fig. S9 |
| | GAGGAGGAGGTTATACACGG | For detection of mutations at At1g70460 | |
| At1g70460-F3 | TCAG | site | |
| | AGTACTGGCCTTCCCTTCCcta | For detection of mutations at At1g70460 | Designated as an filled arrow in Fig. |
| At1g70460-R2 | tc | site and gene cluster inversions | S8A and B |
| At1g70460-R3 | tgcaaaacaaaacaaaacataca | For detection of gene cluster deletions | Designated as R2 in Fig. 3B |
| | -8 | For detection of NHEJ-mediated | |
| At4g16960-F | gtcttgttaggtggtttgatgtta | mutagenesis | |
| | | For detection of mutations at At4g16960 | |
| At4g16960/940-R | CCATTTGATCCAAGTCTTTG | and At4g16940 sites | |
| | | For detection of mutations at At4g16960 | |
| At4g16940-F | agcaccacctcagccccatac | site and gene cluster deletions | Designated as F2 in Fig. 3C |
| | | For detection of mutations at At4g16860 | |
| At4g16860/950-F | tggagggaaggaagacgaagtt | site | |
| | | For detection of mutations at At4g16860 | |
| At4g16860-R | ATTTGTTCCCCTTTCTTGTA | site and gene cluster deletions | |
| At4g16960-F2 | tctgtatcatattagtttagttcg | For detection of gene cluster deletions | Designated as F1 in Fig. 3C |
| At4g16940-R2 | aaagagaataacacagatttattt | For detection of gene cluster deletions | Designated as R2 in Fig. 3C |
| | | For detection of mutations at the ADH1 | _ = ================================== |
| ADH1F | TCGAGGAAGTGGAGGTTGCT | site | |
| | | For detection of mutations at the ADH1 | |
| ADH1R2 | TGGCTGAAGATCAGTCACTCC | site and large chromosomal deletions | Designated as R in Fig. 4A |

Y. Qi *et al.* 11 SI